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Sequence Listing was accepted.
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217-9197 (toll free).
Reviewer: Keisha Douglas
Timestamp: [year=2008; month=11; day=18; hr=15; min=9; sec=15; ms=687; ]
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Application No: 10582868   Version No: 1.1

Input Set:

Output Set:

Started: 2008-11-18 15:06:03.998

Finished: 2008-11-18 15:06:04.601

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 603 ms

Total Warnings: 8

Total Errors: 0

No. of SeqIDs Defined: 15

Actual SeqID Count: 15

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<110>   VIOVY, Jean-Louis
        WEBER, Jeremie
        STOPPA-LYONNET, Dominique

<120>   Method and compositions for assaying mutations in nucleic acids
        and their uses in diagnosis of genetic diseases and cancers

<130>   121697

<140>   10/582,868
<141>   2007-04-25

<150>   PCT/IB03/06016
<151>   2003-12-16

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<170>   PatentIn version 3.5

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<212>   DNA
<213>   Homo Sapiens

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agttaagagc cttgaataat cacaggcaaa tgttgaatga taagaaacaa gctcagatcc   120

agttggaaat taggaaggcc atggaatctg ctgaacaaaa ggaacaaggt ttatcaaggg   180

atgtcacaac cgtgtggaag ttgcgtattg taagctattc aaaaaaagaa aaagattcag   240

gtaagtatgt aaatgctttg tttttatcag ttttattaac ttaaaaaatg accttactaa   300

caaaatgatt a   311

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<212>   DNA
<213>   Homo Sapiens

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ttggtttgaa gaactttctt cagaagctcc accctataat tctgaacctg cagaagaatc   120

tgaacataaa aacaacaatt acgaaccaaa cctatttaaa actccacaaa ggaaaccatc   180

ttataatcag ctggcttcaa ctccaataat attcaaagag caagggctga ctctgccgct   240

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atggtagact ggggagaac                                                       319


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<212>   DNA
<213>   Homo Sapiens

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tctcaagagg agctcattaa ggttgttgat gtggaggagc aacagctgga agagtctggg          180

ccacacgatt tgacggaaac atcttacttg ccaaggcaag atctaggtaa tatttcatct          240

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<212>   DNA
<213>   Homo Sapiens

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gacggaaaca tcttacttgc caaggcaaga tctaggtaat atttcatctg ctgtattgga          180

acaaacactt tgattttact ctgaatccta cataaagata ttctggttaa ccaactttta          240

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<213>   Homo Sapiens

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caacagctgg aagagtctgg gccacacgat ttgacggaaa catcttactt gccaaggcaa          120

gatctaggta atatttcatc tgctgtattg gaacaaacac tttgatttta ctctgaatcc          180

tacataaaga tattctggtt aaccaacttt tagatgtact agtctatcat ggacactttt          240

gttatac                                                                    247


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tgatttgaac accactgaga agcgtgcagc tgagaggcat ccagaaaagt atcagggtag   180

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tatacccatt aatggttcac ctcgaacacc caggcgaggt cagaacagga gtgcacggat   180

agcaaaacaa ctagaaaatg atacaagaat tattgaagtt ctctgtaaag aacatgaatg   240

taatatagat gaggtaattt aacttcatga tttctttaaa acagttaaag tagatttaga   300

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      as identified in Figure 14

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<210> 9
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<220>
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      as identified in Figure 14

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<210>  10
<211>  23
<212>  DNA
<213>  Artificial Sequence

<220>
<223>  amplification primer by PCR of a DNA fragment coding for BRCA-1
       as identified in Figure 14

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<210>  11
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<212>  DNA
<213>  Artificial Sequence

<220>
<223>  amplification primer by PCR of a DNA fragment coding for BRCA-1
       as identified in Figure 14

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<212>  DNA
<213>  Artificial Sequence

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<223>  amplification primer by PCR of a DNA fragment coding for BRCA-1
       as identified in Figure 14

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<210>  13
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<212>  DNA
<213>  Artificial Sequence

<220>
<223>  amplification primer by PCR of a DNA fragment coding for BRCA-1
       as identified in Figure 14

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<210>  14
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<211>    19
<212>    DNA
<213>    Artificial Sequence

<220>
<223>    amplification primer by PCR of a DNA fragment coding for BRCA-1
         as identified in Figure 14

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gggcttaatt aagtataac                                               19


<210>    15
<211>    19
<212>    DNA
<213>    Artificial Sequence

<220>
<223>    amplification primer by PCR of a DNA fragment coding for BRCA-1
         as identified in Figure 14

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